

# Ling-Wen Ding

## List of Publications by Year in descending order

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Version: 2024-02-01

59  
papers

3,797  
citations

136950

32  
h-index

133252

59  
g-index

60  
all docs

60  
docs citations

60  
times ranked

7524  
citing authors

#	ARTICLE	IF	CITATIONS
1	KDM6A Depletion in Breast Epithelial Cells Leads to Reduced Sensitivity to Anticancer Agents and Increased TGF $\beta$ 2 Activity. <i>Molecular Cancer Research</i> , 2022, 20, 637-649.	3.4	4
2	A Transcriptional Regulatory Loop of Master Regulator Transcription Factors, PPARC, and Fatty Acid Synthesis Promotes Esophageal Adenocarcinoma. <i>Cancer Research</i> , 2021, 81, 1216-1229.	0.9	41
3	MNK1 and MNK2 enforce expression of E2F1, FOXM1, and WEE1 to drive soft tissue sarcoma. <i>Oncogene</i> , 2021, 40, 1851-1867.	5.9	11
4	Interplay and cooperation between SREBF1 and master transcription factors regulate lipid metabolism and tumor-promoting pathways in squamous cancer. <i>Nature Communications</i> , 2021, 12, 4362.	12.8	50
5	Repurposing RNA sequencing for discovery of RNA modifications in clinical cohorts. <i>Science Advances</i> , 2021, 7, .	10.3	12
6	RNA-Binding Protein <i>ZFP36L1</i> Suppresses Hypoxia and Cell-Cycle Signaling. <i>Cancer Research</i> , 2020, 80, 219-233.	0.9	40
7	Master transcription factors form interconnected circuitry and orchestrate transcriptional networks in oesophageal adenocarcinoma. <i>Gut</i> , 2020, 69, 630-640.	12.1	68
8	EWS-FLI1 regulates and cooperates with core regulatory circuitry in Ewing sarcoma. <i>Nucleic Acids Research</i> , 2020, 48, 11434-11451.	14.5	18
9	Lineage-Specific Epigenomic and Genomic Activation of Oncogene HNF4A Promotes Gastrointestinal Adenocarcinomas. <i>Cancer Research</i> , 2020, 80, 2722-2736.	0.9	37
10	Integrative Epigenomic Analysis of Transcriptional Regulation of Human CircRNAs. <i>Frontiers in Genetics</i> , 2020, 11, 590672.	2.3	4
11	SOX7 regulates MAPK/ERK-BIM mediated apoptosis in cancer cells. <i>Oncogene</i> , 2019, 38, 6196-6210.	5.9	32
12	LNK suppresses interferon signaling in melanoma. <i>Nature Communications</i> , 2019, 10, 2230.	12.8	21
13	Bromodomain and extraterminal proteins foster the core transcriptional regulatory programs and confer vulnerability in liposarcoma. <i>Nature Communications</i> , 2019, 10, 1353.	12.8	39
14	Mutational and transcriptomic profiling of acute leukemia of ambiguous lineage reveals obscure but clinically important lineage bias. <i>Haematologica</i> , 2019, 104, e200-e203.	3.5	8
15	Clonality and clonal evolution analysis of paediatric ALL based on B-cell receptor/T-cell receptor rearrangement. <i>British Journal of Haematology</i> , 2019, 184, 829-833.	2.5	3
16	Co-targeting poly(ADP-ribose) polymerase (PARP) and histone deacetylase (HDAC) in triple-negative breast cancer: Higher synergism in BRCA mutated cells. <i>Biomedicine and Pharmacotherapy</i> , 2018, 99, 543-551.	5.6	48
17	Super-Enhancer-Driven Long Non-Coding RNA LINC01503, Regulated by TP63, Is Over-Expressed and Oncogenic in Squamous Cell Carcinoma. <i>Gastroenterology</i> , 2018, 154, 2137-2151.e1.	1.3	165
18	Identification of distinct mutational patterns and new driver genes in oesophageal squamous cell carcinomas and adenocarcinomas. <i>Gut</i> , 2018, 67, 1769-1779.	12.1	101

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19	Targeting the vulnerability to NAD <sup>+</sup> depletion in B-cell acute lymphoblastic leukemia. <i>Leukemia</i> , 2018, 32, 616-625.	7.2	29
20	Functional Genome-wide Screening Identifies Targets and Pathways Sensitizing Pancreatic Cancer Cells to Dasatinib. <i>Journal of Cancer</i> , 2018, 9, 4762-4773.	2.5	25
21	Profiling the B/T cell receptor repertoire of lymphocyte derived cell lines. <i>BMC Cancer</i> , 2018, 18, 940.	2.6	10
22	Co-activation of super-enhancer-driven CCAT1 by TP63 and SOX2 promotes squamous cancer progression. <i>Nature Communications</i> , 2018, 9, 3619.	12.8	179
23	Targetable BET proteins- and E2F1-dependent transcriptional program maintains the malignancy of glioblastoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E5086-E5095.	7.1	87
24	The c-MYC-BMI1 axis is essential for SETDB1-mediated breast tumorigenesis. <i>Journal of Pathology</i> , 2018, 246, 89-102.	4.5	28
25	ARID1A and CEBP $\beta$ cooperatively inhibit UCA1 transcription in breast cancer. <i>Oncogene</i> , 2018, 37, 5939-5951.	5.9	24
26	ASXL2 regulates hematopoiesis in mice and its deficiency promotes myeloid expansion. <i>Haematologica</i> , 2018, 103, 1980-1990.	3.5	15
27	Ordering of mutations in acute myeloid leukemia with partial tandem duplication of MLL (MLL-PTD). <i>Leukemia</i> , 2017, 31, 1-10.	7.2	63
28	Genomic and Epigenomic Heterogeneity of Hepatocellular Carcinoma. <i>Cancer Research</i> , 2017, 77, 2255-2265.	0.9	166
29	Mutational profiling of acute lymphoblastic leukemia with testicular relapse. <i>Journal of Hematology and Oncology</i> , 2017, 10, 65.	17.0	16
30	BCL6 promotes glioma and serves as a therapeutic target. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3981-3986.	7.1	58
31	Targeting super-enhancer-associated oncogenes in oesophageal squamous cell carcinoma. <i>Gut</i> , 2017, 66, 1358-1368.	12.1	169
32	Super-Enhancers Promote Transcriptional Dysregulation in Nasopharyngeal Carcinoma. <i>Cancer Research</i> , 2017, 77, 6614-6626.	0.9	103
33	Selinexor (KPT-330) has antitumor activity against anaplastic thyroid carcinoma in vitro and in vivo and enhances sensitivity to doxorubicin. <i>Scientific Reports</i> , 2017, 7, 9749.	3.3	32
34	ZNF750 is a lineage-specific tumour suppressor in squamous cell carcinoma. <i>Oncogene</i> , 2017, 36, 2243-2254.	5.9	90
35	Mutational profiling of a MonoMAC syndrome family with GATA2 deficiency. <i>Leukemia</i> , 2017, 31, 244-245.	7.2	22
36	Diagnosis and relapse: cytogenetically normal acute myelogenous leukemia without FLT3-ITD or MLL-PTD. <i>Leukemia</i> , 2017, 31, 762-766.	7.2	9

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37	Mutational Landscape of Pediatric Acute Lymphoblastic Leukemia. <i>Cancer Research</i> , 2017, 77, 390-400.	0.9	77
38	Comprehensive mutational analysis of primary and relapse acute promyelocytic leukemia. <i>Leukemia</i> , 2016, 30, 1672-1681.	7.2	99
39	Profiling of somatic mutations in acute myeloid leukemia with FLT3-ITD at diagnosis and relapse. <i>Blood</i> , 2015, 126, 2491-2501.	1.4	180
40	Genomic and Functional Analysis of the E3 Ligase PARK2 in Glioma. <i>Cancer Research</i> , 2015, 75, 1815-1827.	0.9	50
41	Activation of protein phosphatase 2A tumor suppressor as potential treatment of pancreatic cancer. <i>Molecular Oncology</i> , 2015, 9, 889-905.	4.6	51
42	LNK (SH2B3): paradoxical effects in ovarian cancer. <i>Oncogene</i> , 2015, 34, 1463-1474.	5.9	21
43	SETDB1 accelerates tumorigenesis by regulating the WNT signalling pathway. <i>Journal of Pathology</i> , 2015, 235, 559-570.	4.5	64
44	Growth inhibition of pancreatic cancer cells by histone deacetylase inhibitor belinostat through suppression of multiple pathways including HIF, NFκB, and mTOR signaling in vitro and in vivo. <i>Molecular Carcinogenesis</i> , 2014, 53, 722-735.	2.7	51
45	Genomic and molecular characterization of esophageal squamous cell carcinoma. <i>Nature Genetics</i> , 2014, 46, 467-473.	21.4	523
46	Laminin-5 $\beta$ -2 (LAMC2) Is Highly Expressed in Anaplastic Thyroid Carcinoma and Is Associated With Tumor Progression, Migration, and Invasion by Modulating Signaling of EGFR. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2014, 99, E62-E72.	3.6	60
47	KPT-330 has antitumour activity against non-small cell lung cancer. <i>British Journal of Cancer</i> , 2014, 111, 281-291.	6.4	59
48	The genomic landscape of nasopharyngeal carcinoma. <i>Nature Genetics</i> , 2014, 46, 866-871.	21.4	317
49	Selective inhibition of unfolded protein response induces apoptosis in pancreatic cancer cells. <i>Oncotarget</i> , 2014, 5, 4881-4894.	1.8	77
50	SOX7 is down-regulated in lung cancer. <i>Journal of Experimental and Clinical Cancer Research</i> , 2013, 32, 17.	8.6	56
51	PIAS4 is an activator of hypoxia signalling via VHL suppression during growth of pancreatic cancer cells. <i>British Journal of Cancer</i> , 2013, 109, 1795-1804.	6.4	32
52	Genomic and functional characterizations of phosphodiesterase subtype 4D in human cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6109-6114.	7.1	59
53	The Characterization of SaPIN2b, a Plant Trichome-Localized Proteinase Inhibitor from <i>Solanum americanum</i> . <i>International Journal of Molecular Sciences</i> , 2012, 13, 15162-15176.	4.1	14
54	Adaptor protein Lnk binds to and inhibits normal and leukemic FLT3. <i>Blood</i> , 2012, 120, 3310-3317.	1.4	38

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55	Improved expression and purification of recombinant human serum albumin from transgenic tobacco suspension culture. <i>Journal of Biotechnology</i> , 2011, 155, 164-172.	3.8	52
56	Culture of <i>Escherichia coli</i> in SOC medium improves the cloning efficiency of toxic protein genes. <i>Analytical Biochemistry</i> , 2009, 394, 144-146.	2.4	12
57	Genome sequence and characterization of a new virus infecting <i>Mikania micrantha</i> H.B.K.. <i>Archives of Virology</i> , 2008, 153, 1765-1770.	2.1	13
58	Using silica particles to isolate total RNA from plant tissues recalcitrant to extraction in guanidine thiocyanate. <i>Analytical Biochemistry</i> , 2008, 374, 426-428.	2.4	56
59	Purification and characterization of native and recombinant SaPIN2a, a plant sieve element-localized proteinase inhibitor. <i>Plant Physiology and Biochemistry</i> , 2007, 45, 757-766.	5.8	8